APPENDIX 1



lalign output for 1 vs. U95825

[ISREC-Server] Date: Thu May 23 5:00:25 MET 2002

resetting	g matrix to	DNA ./ww	wtmp/lalign	/.1485.1.seq	: 5271 nt		
version 1 U95825 scoring	on 2.0uPl	ease cite DNA, gap	l alignmen : Myers an penalties Global ali	d Miller, : -14/-4	CABIOS (19 52 525	989) 4:11-1 71 nt vs. 53 nt	17
	CCGGAGAG CCGGAGAG	:::::::::		::::::::	::::::::::	50 GGGTAGAAATA :::::::::: GGGTAGAAATA 50	::::
	CTGTCATG:::::::CTGTCATG	:::::::	::::::::	::::::::::	: : : : : : : : : : : : : : : : : : :	110 ATATCCGCCTC :::::::: ATATCCGCCTC 110	::::
	TCAAGGAA ::::::: TCAAGGAA	ATATCAGAT ::::::: ATATCAGAT	:::::::: AAAATATCTA	::::::::::	GGTGAGACGA	170 ATTAAAGATGO :::::::::: ATTAAAGATGO 170	::::
./wwwt			200 ATGGACCAGG	210 ACTCTGAAGA	220 AGAAAAGGAG	230 SCTTTATTTA	240 AACC
U95825	:::::: TGAAAACT	::::::: TTTATGGAT	:::::::: ATGGACCAGG	:::::::::		::::::::::::::::::::::::::::::::::::::	::::
./wwwt	TAGCTTTA	CATCTTGCT	:::::::::	:::::::::	::::::	290 AGATGTTCGCT	::::
U95825			TCAGATTTTT 260	TTCTCAAGCA 270	ATCCTGATAAA 280	AGATGTTCGC1 290	7AC 300
./wwwt	TGGTAGCC	TGCTGCCTT	GCTGATATTT			350 AGCTCCTTACA	
U95825	TGGTAGCC	TGCTGCCTT	GCTGATATTT			AGCTCCTTACA 350	
	CCCCTGAT	AAACTAAAG ::::::::	:::::::::	::::::::	::::::::	410 GAAGGGGCTAG SECTION SAAGGGGCTAG	:::
	3	70 30 AGCCCACAA	380 440	390 450 ATTTTTATT1	400 460 CACTTGAGAAC	410 470 CATTGCTTGGG	420 480

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Ū95825	ATACAAAGAGCCCA	CAGTTCAATA	GGTATTTTT	ATTTGCTTGA	GAACATTGCT	TGGGTCA
	430	440	450	460	470	480
	490	500	510	520	530	540
./wwwt	AGTCATATAACATA					
	:::::::::::::::::::::::::::::::::::::::					
U95825	AGTCATATAACATA					0
	490	500	510	520	530	540
	550	560	570	580	590	600
./wwwt	GAACCTTATTTCA					
	:::::::::::::::::::::::::::::::::::::::					
095825	GAACCTTATTTTCA 550	GTTATAAACA 560	ATGGCCACA 570	ATCAGAAAGT(580	CCATATGCAC 590	
	220	200	570	360	390	600
	610	620	630	640	650	660
./wwwt	ACCTTATGAGCTCT		AAGGTGATA	CAGTGTCTCA	GGAGCTTTTG	GATACGG
1105025	ACCTTATGAGCTCT		A A C C T C A T A C	::::::::::::::::::::::::::::::::::::::		:::::::
093023	610	620	630	640	650	660
	670	680	690	700.	710	7.20
./wwwt	TTTTAGTAAATCTG		· ,	raaacaagcaz		
U95825	TTTTAGTAAATCTG					, -
	670	680	690	700	710	720
	720	7.40	5.5 0	7.60	220	=00
/101101101+	730 AGGCTTTACTGAAG	740 AGGACAGCTC	750	760 אכככם עם שם יים	770 רב ככב כת חודים	780 תתתממתר
. / ***********************************	:::::::::::::	::::::::		::::::::	:::::	····
U95825	AGGCTTTACTGAAG					
	730	740	750	760	770	780
	790	800	810	820	830	840
	AGGTTCTGATGCTT	GGGAAAACAT	CTATCAGCGA	ATTTGTCAGA	SCATGTCTTT	
	AGGTTCTGATGCTT	GGGAAAACAT	CTATCAGCGA	ATTTGTCAGA(GCATGTCTTT(GACTTAA
	AGGTTCTGATGCTT::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT	CTATCAGCGA ::::::::: CTATCAGCGA	ATTTGTCAGA(:::::::::::::::::::::::::::::::::::	GCATGTCTTT(:::::::::::::::::::::::::::::::::	GACTTAA :::::: GACTTAA
	AGGTTCTGATGCTT	GGGAAAACAT	CTATCAGCGA	ATTTGTCAGA(GCATGTCTTT(GACTTAA
U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::::::::::::::::::::::::::	CTATCAGCGA ::::::::: CTATCAGCGA 810 870	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880	GCATGTCTTT(::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840
	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA	CTATCAGCGA ::::::::: CTATCAGCGA 810 870 GTCATTTGCT	ATTTGTCAGA(:::::::: ATTTGTCAGA(820 880 GGCTCTCTGT	GCATGTCTTT :::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT
U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::::::::::::::::::::::::::	CTATCAGCGA ::::::::: CTATCAGCGA 810 870 GTCATTTGCT	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880 FGCTCTCTGT	GCATGTCTTT(:::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT
U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::::::::::::::::::::::::::	CTATCAGCGA ::::::::: CTATCAGCGA 810 870 GTCATTTGCT	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880 FGCTCTCTGT	GCATGTCTTT(:::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT
U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA	CTATCAGCGA ::::::::: CTATCAGCGA 810 870 GTCATTTGCT ::::::::: GTCATTTGCT 870	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880 GCTCTCTGTT :::::::::: GCTCTCTCTGTT	GCATGTCTTT(:::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT ::::::: CTTGAAT 900
U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920	CTATCAGCGAELL CTATCAGCGAELO 870 GTCATTTGCTELL CTATTGCTELL CTATTGC	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880 GCTCTCTGTT :::::::::: GCTCTCTGTT 880 940	GCATGTCTTT(:::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT ::::::: CTTGAAT 900 960
U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920	CTATCAGCGA :::::::::: CTATCAGCGA 810 870 GTCATTTGCT ::::::::: GTCATTTGCT 870 930 AGGAGCGCCT	ATTTGTCAGA(::::::::: ATTTGTCAGA(820 880 FGCTCTCTGTT :::::::: FGCTCTCTGTT 880 940 FACAAGTTGTT	GCATGTCTTT(::::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(::::::::: FTTACCCCAG(890 950 FAAACTACTG	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA
U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG AATGATAATG	CTATCAGCGA ::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICAGAGESTICAGAGESTAGAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGAGESTAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	GCATGTCTTT(:::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(:::::::: FTTACCCCAG(890 950 FAAACTACTG(FAAACTACTG(GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA
U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGA(E::::::::: ATTTGTCAGA(820 880 FGCTCTCTGTT E::::::::::::::::::::::::::::::::::	GCATGTCTT(::::::::: GCATGTCTCTCT(830 890 GTTACCCCAG(:::::::::: GTTACCCCAG(890 950 GAAACTACTG(::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA
U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG AATGATAATG	CTATCAGCGA ::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICAGAGESTICAGAGESTAGAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGESTAGAGAGESTAGAGAGESTAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	GCATGTCTTT(:::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(:::::::: FTTACCCCAG(890 950 FAAACTACTG(FAAACTACTG(GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG ::::::::: AATGATAATG 920 980	CTATCAGCGA :::::::::: CTATCAGCGA 810 870 GTCATTTGCT ::::::::: GTCATTTGCT 870 930 AGGAGCGCCT ::::::::: AGGAGCGCCT 930 990	ATTTGTCAGAGESTICAGAGESTICAGAGESTAGAGTTGTTCAGAGTTGTTTCAGAGTTGTTCAGAGTTGTTTCAGAGTTGTTTCAGAGTTGTTTCAGAGTTGTTTCAGAGTTGTTTCAGAGTTGTTTTTAGAGTTGTTTTAGAGTTGTTTTAGAGAGTTGTT	GCATGTCTTT(::::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(::::::::: FTTACCCCAG(890 950 FAAACTACTG(950 1010	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG 920 AATGATAATG 920 980 GATTCAGAAT ::::::::	CTATCAGCGA :::::::::: CTATCAGCGA 810 870 GTCATTTGCT ::::::::: GTCATTTGCT 870 930 AGGAGCGCCT 930 990 TGGCTTCTCAGCAGCGCCAGCCCT ::::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICAGAGESTICAGAGESTAGAGTTGTTAGAGAGTTGTTAGAGAGTTGTTAGAGAGTTGTT	GCATGTCTTT(::::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(::::::::: FTTACCCCAG(890 P50 FAAACTACTG(950 1010 ACTTTGGCAG(::::::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA 960 1020 TGCTACT :::::
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG 920 AATGATAATG 920 980 GATTCAGAAT :::::::: GATTCAGAAT	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGAGES SECTION OF THE PROPERTY OF THE P	GCATGTCTTT S:::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG FAAACTACTG 950 1010 ACTTTGGCAG	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT :::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG 920 AATGATAATG 920 980 GATTCAGAAT ::::::::	CTATCAGCGA :::::::::: CTATCAGCGA 810 870 GTCATTTGCT ::::::::: GTCATTTGCT 870 930 AGGAGCGCCT 930 990 TGGCTTCTCAGCAGCGCCAGCCCT ::::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICAGAGESTICAGAGESTAGAGTTGTTAGAGAGTTGTTAGAGAGTTGTTAGAGAGTTGTT	GCATGTCTTT(::::::::: GCATGTCTCTCT(830 890 FTTACCCCAG(::::::::: FTTACCCCAG(890 P50 FAAACTACTG(950 1010 ACTTTGGCAG(::::::::::::::::::::::::::::::::::::	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA 960 1020 TGCTACT :::::
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA :::::::: AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT :::::::: GATTCAGAAT 980 1040	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SEED SEED SEED SEED SEED SEED SEED SE	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG 950 1010 ACTTTGGCAG ACTTTGGCAG 1010 1070	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT 1020 1080
U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT :::::::: GATTCAGAAT 980 1040 GATATCCATG	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SEED SEED SEED SEED SEED SEED SEED SE	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG PAAACTACTG 950 1010 ACTTTGGCAG 1010 ACTTTGGCAG 1010 1070 TGTGAAATTT	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT 1020 1080 GCTAGCC
U95825 ./wwwt U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT ::::::::: GATTCAGAAT 980 1040 GATATCCATG ::::::::	CTATCAGCGA ::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SEED SEED SEED SEED SEED SEED SEED SE	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG 950 1010 ACTTTGGCAG 1010 1070 TGTGAAATTTC 1070 TGTGAAATTTC	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT 1020 1080 GCTAGCC ::::::
U95825 ./wwwt U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT ::::::::: GATTCAGAAT 980 1040 GATATCCATG ::::::::	CTATCAGCGA ::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SEED SEED SEED SEED SEED SEED SEED SE	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG 950 1010 ACTTTGGCAG 1010 1070 TGTGAAATTTC 1070 TGTGAAATTTC	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT 1020 1080 GCTAGCC ::::::
U95825 ./wwwt U95825 ./wwwt U95825 ./wwwt	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT ::::::::: GATTCAGAAT 980 1040 GATATCCATG ::::::::: GATATCCATG 1040	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SEED OF SEED	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG 950 1010 ACTTTGGCAG 1010 ACTTTGGCAG 1010 1070 TGTGAAATTTC 1070	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 TGCTACT :::::: TGCTACT 1020 1080 GCTAGCC :::::: GCTAGCC 1080
U95825 ./wwwt U95825 ./wwwt U95825 ./wwwt U95825	AGGTTCTGATGCTT ::::::::::::::::::::::::::::::::::	GGGAAAACAT ::::::::: GGGAAAACAT 800 860 AATATTGATA 860 920 AATGATAATG :::::::: AATGATAATG 920 980 GATTCAGAAT :::::::: GATTCCATG :::::::: GATATCCATG 1040 1100	CTATCAGCGA :::::::::::::::::::::::::::::::::	ATTTGTCAGAGESTICS SECTEGATOR SECT	GCATGTCTTT S::::::::: GCATGTCTCT 830 890 TTTACCCCAG 890 950 TAAACTACTG 950 1010 ACTTTGGCAG 1010 1070 TGTGAAATTTC 1070 TGTGAAATTTC 1070 1130	GACTTAA ::::::: GACTTAA 840 900 CTTGAAT :::::: CTTGAAT 900 960 GCAAAAA :::::: GCGAAAA 960 1020 1020 1080 GCTACT :::::: GCTACCC ::::::: GCTAGCC 1080 1140

U95825		:::::::: TCATGAACCA 1090	:::::::: ATCCTGATTTA 1100	:::::::: AGCAAAAGACT 1110	TTAACAGAGTA	:::::::: ATCTTAAAGT0 1130	::::: GAGGT 1140
./wwwt		1150 ACCCTGAGGA	1160 AAGCTATTAGA	1170 ACATGATGTTA	1180 ATTGTGTCAAT	1190 TAGTTACAGC	1200 IGCTA
U95825	CACATG		AAGCTATTAGA 1160				
•		1210	1220	1230	1240	1250	1260
./wwwt			rggtcaatgai :::::::::::			GAGAGAGAAC :::::::::	ATTAG
U95825		ATATTCTTCT 1210	rggtcaatgat 1220	CACTTACTTA 1230	AATTTTGTGG(1240	GAGAGAGAAC 1250	ATTAG 1260
./wwwt	ACAAAC		1280 FACGCAAAGAA				
U95825	ACAAAC		::::::::: TACGCAAAGA/ 1280				
		1330	1340	1350	1360	1370	1380
./wwwt			CAGCTGGAAAA				
U95825			CAGCTGGAAAA 1340				
/www+		1390 TACATATATA	1400 ATTATCAAAA	1410 PAGTATTGATO	1420 SATCGACTAC	1430 FTGTTGAACG	1440 GATCT
	:::::		ATTATCAAAA	:::::::::::	:::::::::	::::::::	:::::
093623	AATIGO	1390	1400	1410	1420	1430	1440
		1450	1460		1480	1490	1500
	::::::		TTCCTCACAAT		::::::::::		::::
U95825	TTGCTC	CAATACATGGT 1450	TTCCTCACAA 1460	TTAGAAACTA 1470	ACAGAACGGA 1480	rgaaatgett 1490	1500
/+	አ ርጥጥርጥ	1510	1520 rggatttaaa:	1530	1540	1550	1560
	:::::	::::::::	:::::::::	:::::::::			:::::
095825	ACTTGT	1510	rggatttaaa: 1520	1530	1540	1550	1560
./wwwt			1580 ATCAAGTAAA				
U95825			:::::::: ATCAAGTAAA 1580				
* .		1630	1640	1650	1660	1670	1680
./wwwt		GTGTCAAGG	CCATATTTTC	AAAAGTGATG	GTTATTACAA	GAAATTTACC	TGATC
U95825			CCATATTTTCA 1640				
./wwwt			1700 ATTTCATGAA				
U95825			::::::::::::::::::::::::::::::::::::::				
		1750	1760	1770	1780	1790	1800

./wwwt	TAAGAAAGCAGTTAGAAGTACTTGTTAGTCCAACATGCTCCTGCAAGCAGGCTGAAGGTT
	:::::::::::::::::::::::::::::::::::::::
U95825	TAAGAAAGCAGTTAGAAGTACTTGTTAGTCCAACATGCTCCTGCAAGCAGGCTGAAGGTT 1750 1760 1770 1780 1790 1800
	1730 1700 1770 1730 1790
	1810 1820 1830 1840 1850 1860
./wwwt	GTGTGCGTGAAATAACTAAGAAGTTGGGCAACCCCAAACAGCCTACAAATCCTTTCCTGG
#10E00E	
095825	GTGTGCGTGAAATAACTAAGAAGTTGGGCAACCCCAAACAGCCTACAAATCCTTTCCTGG 1810 1820 1830 1840 1850 1860
	1010 1020 1030 1040 1000
	1870 1880 1890 1900 1910 1920
./wwwt	AAATGATCAAGTTTCTCTTGGAGAGGATAGCACCTGTGCACATAGATACCGAATCTATCA
1105925	::::::::::::::::::::::::::::::::::::::
093023	1870 1880 1890 1900 1910 1920
	1930 1940 1950 1960 1970 1980
./wwwt	GTGCTCTTATTAAACAAGTGAACAAATCAATAGATGGAACAGCAGATGATGAAGATGAGG
1195825	GTGCTCTTATTAAACAAGTGAACAAATCAATAGATGGAACAGCAGATGATGAAGATGAGG
0,5025	1930 1940 1950 1960 1970 1980
,	1990 2000 2010 2020 2030 2040
./wwwt	GTGTTCCAACTGATCAAGCCATCAGAGCAGGTCTTGAACTGCTTAAGGTACTCTCATTTA
U95825	GTGTTCCAACTGATCAAGCCATCAGAGCAGGTCTTGAACTGCTTAAGGTACTCTCATTTA
	1990 2000 2010 2020 2030 2040
	0050 0050 0070 0000 0000 0100
/1-71-71-7+	2050 2060 2070 2080 2090 2100 CACATCCCATCTCATTCATTCTGCTGAAACATTTGAATCATTACTGGCTTGTCTGAAAA
. / ******	
U95825	CACATCCCATCTCATTCATTCTGCTGAAACATTTGAATCATTACTGGCCTGTCTGAAAA
	2050 2060 2070 2080 2090 2100
	2110 2120 2130 2140 2150 2160
./wwwt	TGGATGATGAAAAAGTAGCAGAAGCTGCACTACAAATTTTCAAAAACACAGGAAGCAAAA
U95825	TGGATGATGAAAAAGTAGCAGAAGCTGCACTACAAATTTTCAAAAAACACAGGAAGCAAAA 2110 2120 2130 2140 2150 2160
	2110 2120 2130 2140 2150 2160
	2170 2180 2190 2200 2210 2220
./wwwt	$\tt TTGAAGAGGATTTTCCACACATCAGATCAGCCTTGCTTCCTGTTTTACATCACAAATCTA$
1105005	TTGAAGAGGATTTTCCACACATCAGATCAGCCTTGCTTCCTGTTTTACATCACAAATCTA
095625	2170 2180 2190 2200 2210 2220
	2230 2240 2250 2260 2270 2280
./wwwt	AAAAAGGACCCCCCGTCAAGCCAAATATGCCATTCATTGTATCCATGCGATATTTTCTA
U95825	AAAAAGGACCCCCCGTCAAGCCAAATATGCCATTCATTGTATCCATGCGATATTTTCTA
	2230 2240 2250 2260 2270 2280
/r.m.m.a+	2290 2300 2310 2320 2330 2340 GTAAAGAGACCCAGTTTGCACAGATATTTGAGCCTCTGCATAAGAGCCTAGATCCAAGCA
./www.c	GIAAAGAGACCCAGIIIGCACAGAIAIIIGAGCCICIGCAIAAGAGCCIAGAICCAAGCA
U95825	GTAAAGAGGCCCAGTTTGCACAGATATTTGAGCCTCTGCATAAGAGCCTAGATCCAAGCA
	2290 2300 2310 2320 2330 2340
	2350 2360 2370 2380 2390 2400
./wwwt	ACCTGGAACATCTCATAACACCATTGGTTACTATTGGTCATATTGCTCTCCTTGCACCTG
U95825	ACCTGGAACATCTCATAACACCATTGGTTACTATTGGTCATATTGCTCTCCTTGCACCTG
	2350 2360 2370 2380 2390 2400

./wwwt	2410 ATCAATTTGCTGCTC					
U95825	ATCAATTTGCTGCTC					
./wwwt	2470 TGAATGATCGGCTT					
U95825	TGAATGATCGCCTTC					
./wwwt	2530 CTCCTGAGACAATG					
U95825	CTCCTGAGACAATGC 2530					
./wwwt	2590 TGAAAAATAATCACA	2600 AGTAAATCAG	2610 GAACTTCTAG	2620 CCTTAAGATT	2630 GCTAACAACA	2640 ATATTGC
U95825	TGAAAAATAATCACA 2590				GCTAACAACA	
./wwwt	2650 ATAGTGATGGAGAC					
U95825	ATAGTGATGGAGAC					
./wwwt	2710 GACTTGCTGCTGGGA					
U95825	GACTTGCTGCTGGGA					
./wwwt	2770 TCACATTAGAACAA					
U95825	TCACATTAGAACAA 2770					
./wwwt	2830 GACAAGTGTTTGCC	2840 CAGAAACTTC	2850 ACAAAGGCC	2860 FTTCCCGTTT	2870 ACGGCTTCCA	2880 CTTGAGT
U95825	GACAAGTGTTTGCCC					
./wwwt	2890 ATATGGCAATCTGT0	2900 GCCCTTTGTG	2910 CAAAAGATCO	2920 CTGTAAAGGA	2930 GAGAAGAGCT	2940 CATGCTA
U95825	ATATGGCAATCTGTC 2890					
./wwwt	2950 GGCAATGTTTGGTG	2960 AAAAATATAA	2970 ATGTAAGGC0	2980 GGAGTATCT	2990 GAAGCAGCAT	3000 GCAGCTG
U95825	GGCAATGTTTGGTG					
./wwwt	3010 TTAGTGAAAAATTA	3020 FTGTCTCTTC	3030 TACCAGAGTA	3040 ATGTTGTTCC	3050 ATATACAATT	3060 CACCTTT
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./wwwt	3070 3080 3090 3100 3110 3120 TGGCACATGACCCAGATTATGTCAAAGTACAGGATATTGAACAACTTAAAGATGTTAAAG
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./wwwt	3130 3140 3150 3160 3170 3180 AATGTCTTTGGTTTGTTCTGGAAATATTAATGGCTAAAAATGAAAATAACAGTCACGCTT
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./wwwt	3190 3200 3210 3220 3230 3240 TTATCAGAAAGATGGTAGAAAATATTAAACAAACAAAAGATGCCCAAGGACCAGATGATG
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./wwwt	3310 3320 3330 3340 3350 3360 AGAGTACTACAGTTTGGAATCTCCTAAAGACCCGGTACTACCAGCTCGTTTCTTCA
U95825	AGAGTACTACATACAGTTTGGAATCTCCTAAAGACCCGGTACTACCAGCTCGTTTCTTCA 3310 3320 3330 3340 3350 3360
./wwwt	3370 3380 3390 3400 3410 3420 CTCAACCTGACAAGAATTTCAGTAACACCAAAAATTATCTGCCTCCTGAAATGAAATCAT
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./wwwt	3430 3440 3450 3460 3470 3480 TTTTCACTCCTGGAAAACCTAAAACAACCAATGTTCTAGGAGCTGTTAACAAGCCACTTT
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./wwwt	3490 3500 3510 3520 3530 3540 CATCAGCAAGCAAGCAATCATCACGAATGGAAACTGTAAGCAATGCAA
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./wwwt	3550 3560 3570 3580 3590 3600 GCAGCAGCTCAAATCCAAGCTCTCCTGGAAGAATAAAGGGGAGGCTTGATAGTTCTGAAA
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./wwwt	3610 3620 3630 3640 3650 3660 TGGATCACAGTGAAAATGAAGATTACACAATGTCTTCACCTTTGCCGGGGAAAAAAAGTG
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./wwwt	3670 3680 3690 3700 3710 3720 ACAAGAGAGACGACTCTGATCTTGTAAGGTCTGAATTGGAGAAGCCTAGAGGCAGGAAAA
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	3730	3740	3750	3760	3770	3780
./wwwt	AAACGCCCGTCACA		GAGAAATTAGO	TATGGATGA	CTTGACTAAG	TTGGTAC
U95825	AAACGCCCGTCACA			TATGGATGA	CTTGACTAAG	
	3730	3740	3750	3760	3770	3780
	3790	3800	3810	3820	3830	3840
./wwwt	AGGAACAGAAACCT					
1195825	::::::::::::::::::::::::::::::::::::::					
033023	3790	3800	3810	3820	3830	3840
	3850	3860	3870	3880	3890	3900
./wwwt	CTGATGAACAGCAG					
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. /wwwt	3910 ATGAACAGAATAGT	3920 CCGCCAAAA	3930 AAGGGTAAAA	3940 BAGGCCGACC	3950 ACCAAAACCT	3960 Сттсстс
	:::::::::::::::::::::::::::::::::::::::		: : : : : : : : : :	:::::::		::::::
U95825	ATGAACAGAATAGT 3910	CCGCCAAAAI 3920	AAGGGTAAAAA 3930	SAGGCCGACC 3940	ACCAAAACCT 3950	CTTGGTG 3960
	3910	3920	3930	3940	3930	3960
	3970	3980	3990	4000	4010	4020
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	4030	4040	4050	4060	4070	4080
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	4090	4100	4110	4120	4130	4140
./wwwt	AGAAGTCCAAAAGC					
1105025	::::::::::::::::::::::::::::::::::::::					
093023	4090	4100	4110	4120	4130	4140
	4150	41.00	4170	4100	4100	4200
./wwwt	4150 CTGAATCTAGTGCA	4160 ATTGAATCC	4170 ACACAGTCCAC	4180 CACCACAGAA	4190 AGGACGAGGA	
	:::::::::::::::::::::::::::::::::::::::					
U95825	CTGAATCTAGTGCA 4150	ATTGAATCC 4160	ACACAGTCCAC 4170	ACCACAGAA 4180	AGGACGAGGA 4190	AGACCAT 4200
	4130			1100	4150	
/+	4210 CAAAAACGCCATCA	4220	4230	4240	4250	4260
./www.	CAAAAACGCCATCA				-	-
U95825	CAAAAACGCCATCA					
	4210	4220	4230	4240	4250	4260
,	4270	4280	4290	4300	4310	4320
./wwwt	AAACCAATTTCAAA					
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	4270	4280	4290	4300	4310	4320
	4330	4340	4350	4360	4370	4380
./wwwt	TTTAAATTCCATAT	ATTTAGCCC	CATTACACTAC	GTACGGCGG	CGAAGTGCTA	AAAGGGA
	:::::::::::::::::::::::::::::::::::::::	:::::::::	::::::::::::	:::::::	:::::::::	::::::

U95825	TTTAAATTCCATATA 4330	TTTAGCCCCA 4340	TTACACTAGO 4350	TACGGCGGC 4360	GAAGTGCTAA 4370	AAAGGGA 4380
./wwwt	4390 ACGGCGATGAACAAA	4400 TGTAATTAAT	4410 AACTTTCTCT	4420 TGTGAAAGCT	4430 TTGGAAAAA	4440 CTTTTT
·		:::::::::	:::::::::		::::::::::	::::::
U95825	ACGGCGATGAACAAA	TGTAATTAAT	AACTTTCTCI	GTGAAAGCT	TTGGAAAAA	CTTTTT
	4390	4400	4410	4420	4430	4440
	4450	4460	4470	4480	4490	4500
./wwwt	TTTTTTTTTTTTT					
		:::::::::::::::::::::::::::::::::::::::				
095825	TTTTTTTTTTTTTTT 4450	TTGGTCAAGC 4460	TTGAGGCTGA 4470	AATAAAGCCT 4480	TTGATGCACA 4490	4500
. ,	4510		4530	4540	4550	4560
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033023	4510	4520	4530	4540	4550	4560
	4570	4580	4590	4600	4610	4620
./wwwt	TGCTTTAGCCATACA					
•,	:::::::::::::::::::::::::::::::::::::::					
U95825	TGCTTTAGCCATACA					
•	4570	4580	4590	4600	4610	4620
	4630	4640	4650	4660	4670	4680
./wwwt	TGGCTATGTAGACAT					
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	4690	4700	4710	4720	4730	4740
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	:::::::::::::::::::::::::::::::::::::::					
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	4690	4700	4710	4720	47.30	4740
	4750	4760	4770	4780	4790	4800
./wwwt	TTTATTGGATCTATT					

095825	TTTATTGGATCTATT 4750	GATTTGAAAA 4760	GAATTTGTT <i>E</i> 4770	AGGATAGATC 4780	11AAGCAGTA 4790	4800
	4/30	4700	4770	4700	4730	4000
	4810	4820	4830	4840	4850	4860
./wwwt	CAGTGTTTGTATTTG					

095825	CAGTGTTTGTATTTG	4820	AATTTTACTG	FIGAAAAAAAA 4840	4850	4860
	4810	4820	4030	4040	4030	4000
	4870	4880	4890	4900	4910	4920
./wwwt	TGGTGTCATTTTCTT					
	:::::::::::::::::::::::::::::::::::::::					
095825	TGGTGTCATTTTCTT					
	4870	4880	4890	4900	4910	4920
	4930	4940	4950	4960	4970	4980
./wwwt	TATCTTACCTAGTGT	TTACTCCTGG	GCACCCTTA	ATCTTCAGAG	GTGCTAAATT	TGTCTGC
	:::::::::::::::::::::::::::::::::::::::					
U95825	TATCTTACCTAGTGT					
	4930	4940	4950	4960	4970	4980
	4000	5000	E 0 1 0	5020	E030	E 0 4 0
	4990	3000	5010	3020	5030	5040

U95825	CATTACACCAGAAGG	:::::::: ATGCCTCTGA 5000	TAGGAGGACA 5010	:::::::: ACCATGCAAA 5020	::::::: TTGTGAAATA(5030	::::: GTCCT 5040
	5050		5070	5080	5090	5100
./wwwt	GAAGTTCTTGGATTA	CTTTACACCT	CAGTATTGAT'	TTGTCCCAGA.	ATTTTCTGGC	CTTTC
		:::::::::	:::::::::		:::::::::	::::
U95825	GAAGTTCTTGGATTA					
	5050	5060	5070	5080	5090	5100
	5110		5130			5160
./wwwt	ATGGCAATGAAAATT	TTAAGAAGAA.	AGATTTAAAG'	TTTAATTT	TTAAAGAGTG'	rgtta
	:::::::::::::::::::::::::::::::::::::::	:::::::::	::::::::	::::::::	:::::::::	:::::
U95825	ATGGCAATGAAAATT					IGTTA
	5110	5120	5130	5140	5150	5160
	5170		5190	5200		5220
./wwwt	TAAAATAATGTACTG	AATTCTTTAT	CCCATTTTAT	CATCCTTTCA	GTTTTTATTA	ATCTA
	:::::::::::::::::::::::::::::::::::::::	:::::::::	::::::::	::::::::::	:::::::::	:::::
U95825	TAAAATAATGTACTG					
	5170	5180	5190	5200	5210	5220
				•		
	5230		5250	5260	5270	
./wwwt	CTGTATCAATAAAAT	TCTGTAATTT	GAATGAGTAA	AAAAAAAAA	AAAAA	
	:::::::::::::::::::::::::::::::::::::::	:::::::::::::::::::::::::::::::::::::::	:::::::			
U95825	CTGTATCAATAAAAT'					
	5230	5240	5250			

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